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LAMPIRAN

1. Gambar Pengambilan Sampel

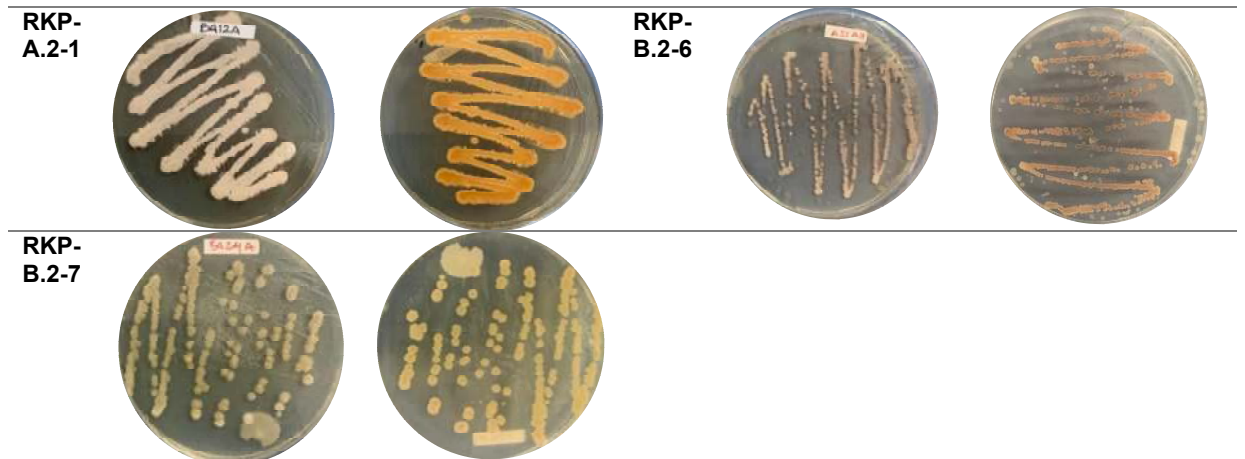


2. Isolasi Actinomycetes



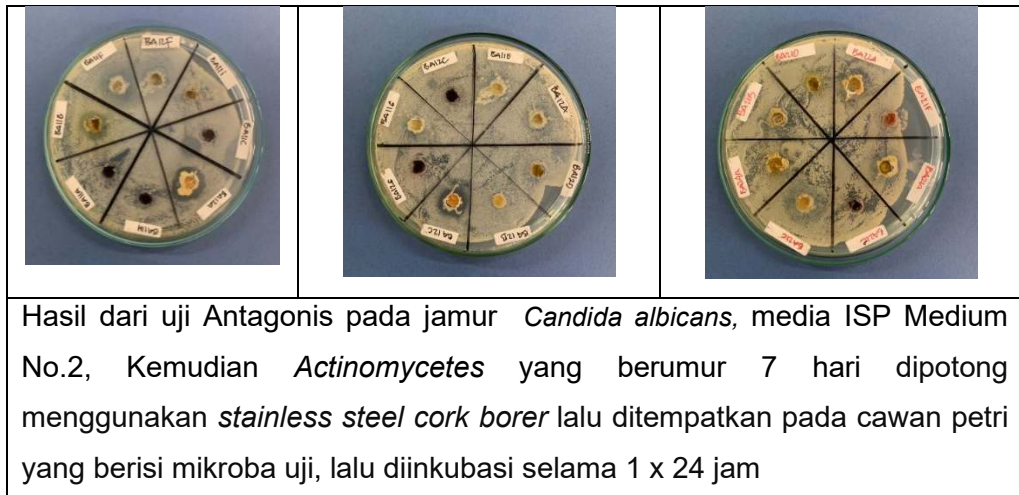
3. Hasil Dari Pemurnian *Actinomycetes*

Kode isolat	Ariel Mycelium	Substrater Mycelium	Kode isolat	Ariel Mycelium	Substrater Mycelium
RKP-A.1-2			RKP-A.2-6		
RKP-A.1-3			RKP-A.2-7		
RKP-A.1-5			RKP-B.1-1		
RKP-A.1.6			RKP-B.2-1		
RKP-A.1-7			RKP-B.2-2		
RKP-A.1-8			RKP-B.2-3		
RKP-A.1-9			RKP-B.2.4		

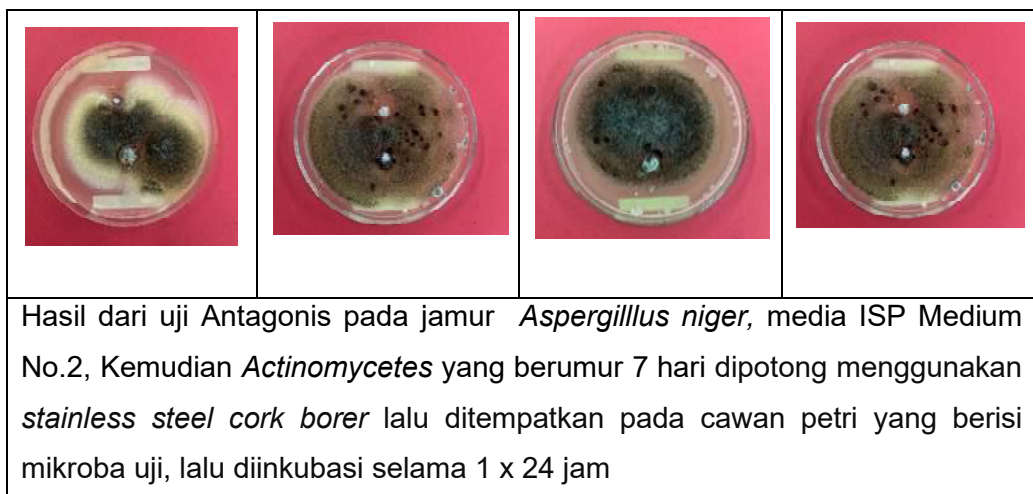


4. Gambar hasil uji antagonis

a. *Candida albicans*



b. *Aspergillus niger*

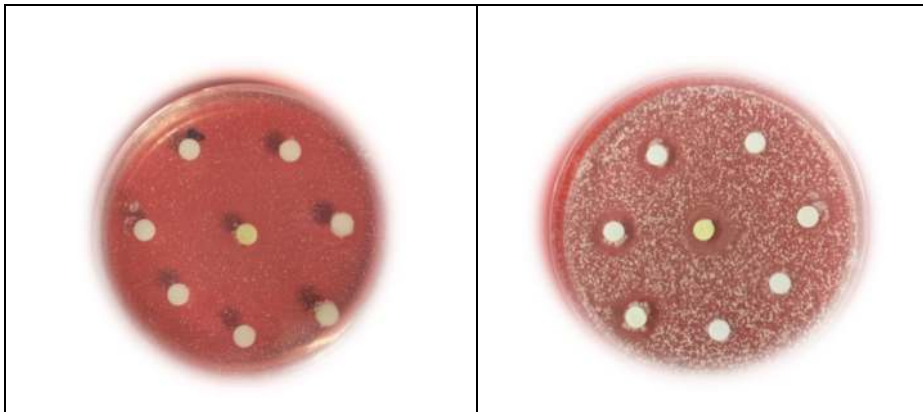


5. Fermentasi



Fermentasi dilakukan dalam kondisi tershaker dengan kecepatan 150 rpm.

6. Uji Aktivitas

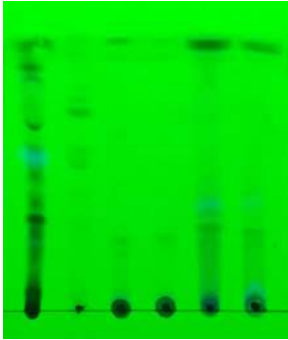
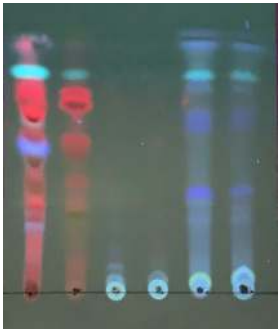



Hasil pengujian aktivitas jamur *Candida albicans*

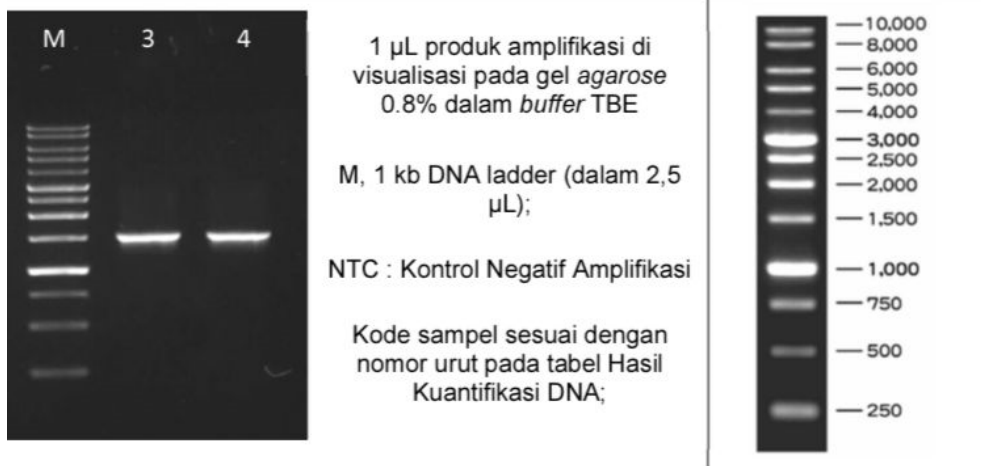
7. Kromatografi Lapis Tipis

	
Maserasi tumbuhan paku dengan menggunakan pelarut etil asetat	Hasil dari maserasi yang telah disaring menjadi ekstrak

Hasil KLT

		
UV 366nm	UV 254 nm	H ₂ SO ₄ 10% Setelah dipanaskan

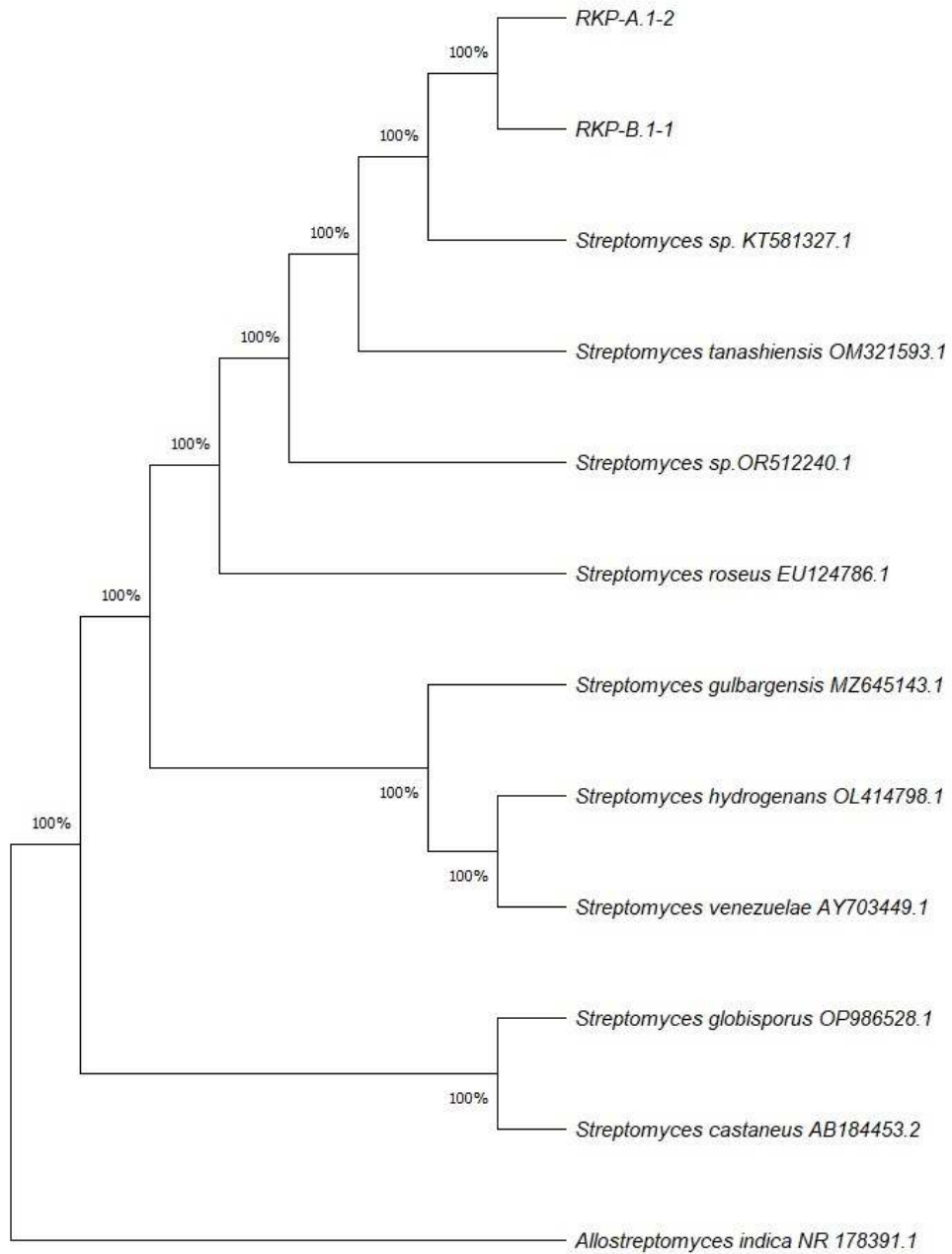
8. Identifikasi Molekuler
a. Elektroforesis Produk PCR



b. Hasil Sequensing – Produk Amplifikasi

NO	Kode Sampel	Sekuens		
1	RKP-A.1-1	<i>Sequence Assembly 1390bp</i>		
		<pre> 1 TGCAGTCGAA CGATGAAGCC CTTCCGGGTG GATTAGTGGC GAACGGGTGA GTAACACGTG 61 GGC AATCTGC CCTTCACTCT GGGACAAGCC CTGGAAACGG GGTCTAATAC CGGATACGAG 121 TCTGGGAGGC ATCTCCTGGA CTGGAAAGCT CCGGCGGTGA AGGATGAGCC CGCGGCCTAT 181 CAGCTTGTG GTGGGGTAAT GGCCTACCAA GGCACGACG GGTAGCCGGC CTGAGAGGGC 241 GACCGGCCAC ACTGGGACTG AGACACGGCC CAGACTCCTA CGGGAGGCAG CAGTGGGGAA 301 TATFGCACAA TGGGC GAAAG CCTGATGCAG CGACGCCGCG TGAGGGATGA CGGCCCTCCG 361 GTTGTA AAC TCTTTCAGCA GGAAGAAGC GAAAGTGACG GTACCTGCAG AAGAAGCGCC 421 GGCTAAC TAC GTGCCAGCAG CCGCGGTAAT ACGTAGGGCG CAAGCGTTGT CCGGAATTAT 481 TGGGCGTAAA GAGCTCGTAG GCGGCTTGTG ACGTCCGGTG TGAAGCCCG GGGCTTAACC 541 CCGGCTCTGC ATCCGATACG GGCAGGCTAG AGTGTGGTAG GGGAGATCGG AATTCCTGGT 601 GTAGCGGTGA AATGCGCAGA TATCAGGAGG AACACCGGTG GCGAAGCGCG ATCTCTGGGC 661 CAT TACTGAC GCTGAGGAGC GAAAGCGTGG GGAGCGAACA GGATTAGATA CCTTGGTAGT 721 CCACGCCGTA AACGTTGGGA ACTAGGTGTT GCGACATTC CACGTCTCGC GTGCCGAGC 781 TAACGCATTA AGTTC CCGC CTGGGGAGTA CCGCCGCAAG GCTAAAACCT AAAGGAATTG 841 ACGGGGGCCC GCACAAGCAG CCGAGCATGT GGCTTAATTC GACGCAACGC GAAGAACCCT 901 ACCAAGGCTT GACATAFACC GGAAGGCATT AGAGATAGTG CCCCCTTGT GGTCCGGTAT 961 CAGGTGGTGC ATGGCTGTG TCAGCTCGTG TCGTGAGATG TTGGGTTAAG TCCCGCAACG 1021 AGCGCAACCC TTGTCCTGTG TTGCCAGCAT GCCCTTCGGG GTGATGGGGA CTCACAGGAG 1081 ACCGCCGGGG TCAACTCGGA GGAAGGTGGG GACGACGTCA AGTCATCATG CCCCTTATGT 1141 CTTGGGCTGC ACACGTGTGA CAATGGCCGG TACAAGAGC TCCGATCCCG TGAGGGCGGAG 1201 CGAATCTCAA AAAGCCGGTC TCAGTTCGGA TTGGGGTCTG CAACTCGACC CCATGAAGTC 1261 GGAGTTGCTA GTAATCGCAG ATCAGCATTG CTGCGGTGAA TACGTTC CCGGCTTGTAC 1321 ACACCGCCCG TCACGTCACG AAAGTCCGTA ACACCCGAAG CCGGTGGCCC AACCCCTTGT 1381 GGGAGGGAGC </pre>		
		2	RKP-B.1-1	<i>Sequence Assembly 1383bp</i>
				<pre> 1 GCAGTCGAAC GATGAAGCCC TTCGGGGTGG ATTAGTGGCG AACGGGTGAG TAACACGTGG 51 GCAATCTGCC CTTCACTCTG GGACAAGCCC TGGAAACGGG GTCTAATACC GGATACGAGT 121 CTGGGAGGCA TCTCCTGGAC TGGAAGGCTC CCGGCGGTGAA GGATGAGCCC GCGGCCTATC 181 AGCTTGTGTTG TGGGGTAATG GCCTACCAAG GCGACGACGG GTAGCCGGCC TGAGAGGGCG 241 ACCGGCCACA CTGGGACTGA GACACGGCCC AGACTCCTAC GGGAGGCAGC AGTGGGGAAT 301 ATTGCACAAT GGGCGAAAGC CTGATGCAGC GACGCCGCGT GAGGGATGAC GGCCTTCGGG 361 TTGTAAACCT CTTTCAGCAG GGAAGAAGCG AAAGTGACGG TAGCTGCAGA AGAAGCGCCG 421 GCTAACTACG TGCCAGCAGC CGCGGTAATA CGTAGGGCGC AAGCGTTGTC CGGAATTATT 481 GGGCGTAAAG AGCTCGTAGG CCGCTTGTCA CGTCGGGTGT GAAAGCCCGG GGCTTAACCC 541 CGGGTCTGCA TCCGATACGG GCAGGCTAGA GTGTGGTAGG GGAGATCGGA ATTCCTGGTG 501 TAGCGGTGAA ATGCGCAGAT ATCAGGAGGA ACACCGGTGG CGAAGGCGGA TCTCTGGGCC 561 ATTACTGACG CTGAGGAGCG AAAGCGTGGG GAGCGAACAG GATTAGATAC CCTGGTAGTC 721 CACGCCGTAA ACGTTGGGAA CTAGGTGTTG GCGACATTCC ACGTCTCGG TGCCCGAGCT 781 AACGCATTAA GTTCCCGGCC TGGGGAGTAC GGCCGCAAGG CTAAAAC TCA AAGGAATTGA 341 CGGGGGCCCG CACAAGCAGC GGAGCATGTG GCTTAATTCG ACGCAACCGG AAGAACCCTA 301 CCAAGGCTTG ACATATACCG GAAAGCATTG GAGATAGTGC CCCCTTGTG GTCGGTATAC 961 AAGTGGTGCA TGGCTGTCGT CAGCTCGTGT CGTGAGATGT TGGGTTAAGT CCCGCAACGA 1021 GCGCAACCCT TGTCCTGTGT TGCCAGCATG CCCTTCGGGG TGATGGGGAC TCACAGGAGA 1081 CCGCCGGGGT CAACTCGGAG GAAAGTGGGG ACGACGTC AATCATCTGC CCCTTATGTC 1141 TTGGGCTGCA CACGTGCTAC AATGGCCGGT ACAAAGAGCT GCGATGCCGT GAGGCGGAGC 1201 GAATCTCAA AAGCCGGTCT CAGTTCGGAT TGGGGTCTGC AACTCGACCC CATGAAGTCG 1261 GAGTTGCTAG TAATCGCAGA TCAGCATTGC TGCGGTGAAT ACGTTC CCGGCTTGTACA 1321 CACCCGCCGT CACGTCACGA AAGTCCGTTA CACCCGAAGC CCGTGGCCCA ACCCCCTTGT 1381 GGA </pre>

Pohon Filogenik



C. Hasil Top 10 Hit BLAST terhadap NCBI, Excluding Uncultured

Kode Sampel	Tautan Hasil																																																																													
<p>G-3231-3 RKP-A.1-2</p>	<p>Sample Sequences</p> <table border="1"> <thead> <tr> <th>Description</th> <th>Max Score</th> <th>Total Score</th> <th>Query Cover</th> <th>E value</th> <th>Per. Ident</th> <th>Accession</th> </tr> </thead> <tbody> <tr> <td>Streptomyces tanashiensis strain WMF812-qm 16S ribosomal RNA gene, partial sequence</td> <td>2567</td> <td>2567</td> <td>100%</td> <td>0.0</td> <td>100.00%</td> <td>OM321593.1</td> </tr> <tr> <td>Streptomyces tanashiensis strain WR78-qm 16S ribosomal RNA gene, partial sequence</td> <td>2567</td> <td>2567</td> <td>100%</td> <td>0.0</td> <td>100.00%</td> <td>OM320200.1</td> </tr> <tr> <td>Streptomyces sp. XT-11 16S ribosomal RNA gene, partial sequence</td> <td>2562</td> <td>2562</td> <td>100%</td> <td>0.0</td> <td>99.93%</td> <td>KT581327.1</td> </tr> <tr> <td>Streptomyces sp. CC5 16S ribosomal RNA gene, partial sequence</td> <td>2562</td> <td>2562</td> <td>100%</td> <td>0.0</td> <td>99.93%</td> <td>KF815090.1</td> </tr> <tr> <td>Streptomyces tanashiensis strain MJM10101 16S ribosomal RNA gene, partial sequence</td> <td>2562</td> <td>2562</td> <td>100%</td> <td>0.0</td> <td>99.93%</td> <td>GU350490.1</td> </tr> <tr> <td>Streptomyces sp. XAS585 16S ribosomal RNA gene, partial sequence</td> <td>2562</td> <td>2562</td> <td>100%</td> <td>0.0</td> <td>99.93%</td> <td>GQ395240.1</td> </tr> <tr> <td>Streptomyces tanashiensis strain HBUM174077 16S ribosomal RNA gene, partial sequence</td> <td>2562</td> <td>2562</td> <td>100%</td> <td>0.0</td> <td>99.93%</td> <td>FJ486422.1</td> </tr> <tr> <td>Streptomyces sp. strain F-29 16S ribosomal RNA gene, partial sequence</td> <td>2556</td> <td>2556</td> <td>100%</td> <td>0.0</td> <td>99.86%</td> <td>MG266317.1</td> </tr> <tr> <td>Streptomyces sp. strain F-28 16S ribosomal RNA gene, partial sequence</td> <td>2556</td> <td>2556</td> <td>100%</td> <td>0.0</td> <td>99.86%</td> <td>MG266316.1</td> </tr> <tr> <td>Streptomyces sp. strain Sed7v 16S ribosomal RNA gene, partial sequence</td> <td>2556</td> <td>2556</td> <td>100%</td> <td>0.0</td> <td>99.86%</td> <td>OR512240.1</td> </tr> </tbody> </table> <p>https://www.ncbi.nlm.nih.gov/nuccore/OM321593.1,OM320200.1,KT581327.1,KF815090.1,GU350490.1,GQ395240.1,FJ486422.1,MG266317.1,MG266316.1,OR512240.1</p>	Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession	Streptomyces tanashiensis strain WMF812-qm 16S ribosomal RNA gene, partial sequence	2567	2567	100%	0.0	100.00%	OM321593.1	Streptomyces tanashiensis strain WR78-qm 16S ribosomal RNA gene, partial sequence	2567	2567	100%	0.0	100.00%	OM320200.1	Streptomyces sp. XT-11 16S ribosomal RNA gene, partial sequence	2562	2562	100%	0.0	99.93%	KT581327.1	Streptomyces sp. CC5 16S ribosomal RNA gene, partial sequence	2562	2562	100%	0.0	99.93%	KF815090.1	Streptomyces tanashiensis strain MJM10101 16S ribosomal RNA gene, partial sequence	2562	2562	100%	0.0	99.93%	GU350490.1	Streptomyces sp. XAS585 16S ribosomal RNA gene, partial sequence	2562	2562	100%	0.0	99.93%	GQ395240.1	Streptomyces tanashiensis strain HBUM174077 16S ribosomal RNA gene, partial sequence	2562	2562	100%	0.0	99.93%	FJ486422.1	Streptomyces sp. strain F-29 16S ribosomal RNA gene, partial sequence	2556	2556	100%	0.0	99.86%	MG266317.1	Streptomyces sp. strain F-28 16S ribosomal RNA gene, partial sequence	2556	2556	100%	0.0	99.86%	MG266316.1	Streptomyces sp. strain Sed7v 16S ribosomal RNA gene, partial sequence	2556	2556	100%	0.0	99.86%	OR512240.1
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9. Hasil Determinasi Tanaman



LABORATORIUM BOTANI DEPARTEMEN BIOLOGI
FAKULTAS MATEMATIKA DAN ILMU PENGETAHUAN ALAM
UNIVERSITAS HASANUDDIN, KAMPUS TAMALANREA
JL. PERINTIS KEMERDEKAAN KM. 10 TLP. (0411) 585466, Fax: 620411 MAKASSAR 90915

Nomor : 061/UN4.11.9/BIO-BOT/PL-03/2024
Lampiran : -
Hal : Hasil Identifikasi dan Determinasi Tanaman

Kepada Yth,
Maulidiah Alda Sami
Di-

Tempat

Dengan hormat,

Bersama ini, kami sampaikan hasil identifikasi dan determinasi tanaman *Nephrolepis cordifolia* (L.) C. Presl. yang saudara (i) kirimkan. Identifikasi dilakukan oleh staf peneliti Laboratorium Botani Departemen Biologi FMIPA Unhas dengan hasil sebagai berikut:

Regnum : Plantae
Divisio : Pteridophyta
Classis : Filicinae
Sub Classis : Leptosporangiateae (Filices)
Ordo : Polypodiales
Familia : Polypodiaceae
Genus : *Nephrolepis*
Species : *Nephrolepis cordifolia* (L.) C. Presl.
Sinonim : *Aspidium cordifolium* (L.) Sw., *Polypodium cordifolium* L.
Nama Lokal : Paku sepat, Pakis kelabang (Indonesia), Narrow swordfern (Inggris)

Kunci Determinasi:

1a- (Golongan 1. Paku dan paku-pakuan)
17b-18b-19b-22b-23b-24b-25b-26b- (Fam 11. Polypodiaceae)
1b-5b-10a-11a- (9. *Nephrolepis*)
1a- (*Nephrolepis cordifolia*)

Buku Acuan:

1. Gembong Tjitrosoepomo. 2011. Taksonomi Tumbuhan (Schizophyta, Thallophyta, Bryophyta, Pteridophyta).
2. Dr. c. g. j. Van Steenis, dkk. 2013. FLORA.
3. *Nephrolepis cordifolia* (L.) C. Presl | Plants of the World Online | Kew Science

Demikian hasil identifikasi kami untuk diketahui dan dipergunakan sebagaimana mestinya.

Makassar, 28 Juni 2024

Kepala Laboratorium

Dr. Andi Ilham Latunra, M.Si
NIP.19670207 199103 1 001